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1600

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/155,327F

DATE: 01/21/2003

TIME: 12:26:16

Input Set : N:\Crf4\01172003\I155327G.raw

Output Set: N:\CRF4\01212003\I155327F.raw

1 <110> APPLICANT: AMRAD Operations Pty Ltd  
 2 <120> TITLE OF INVENTION: A NOVEL MAMMALIAN GENE, bcl-2, BELONGS TO THE bcl-2  
 3 FAMILY OF APOPTOSIS-CONTROLLING GENES  
 4 <130> FILE REFERENCE: 2096584  
 C--> 5 <140> CURRENT APPLICATION NUMBER: US/09/155,327F  
 C--> 6 <141> CURRENT FILING DATE: 1996-03-27  
 7 <150> PRIOR APPLICATION NUMBER: PN8965  
 8 <151> PRIOR FILING DATE: 1996-03-27  
 9 <160> NUMBER OF SEQ ID NOS: 15  
 10 <170> SOFTWARE: PatentIn Ver. 2.1  
 12 <210> SEQ ID NO: 1  
 13 <211> LENGTH: 33  
 14 <212> TYPE: DNA  
 15 <213> ORGANISM: Mouse  
 16 <220> FEATURE:  
 17 <221> NAME/KEY: modified\_base  
 18 <222> LOCATION: 16  
 19 <223> OTHER INFORMATION: n is inosine  
 20 <220> FEATURE:  
 21 <221> NAME/KEY: modified\_base  
 22 <222> LOCATION: 19  
 23 <223> OTHER INFORMATION: n is inosine  
 24 <220> FEATURE:  
 25 <221> NAME/KEY: modified\_base .  
 26 <222> LOCATION: 22  
 27 <223> OTHER INFORMATION: n is inosine  
 28 <220> FEATURE:  
 29 <221> NAME/KEY: modified\_base  
 30 <222> LOCATION: 25  
 31 <223> OTHER INFORMATION: n is inosine  
 32 <400> SEQUENCE: 1

33

W--> 33 gctctagaac tgggnhgnr tngtngcatt ytt  
 35 <210> SEQ ID NO: 2  
 36 <211> LENGTH: 9  
 37 <212> TYPE: PRT  
 38 <213> ORGANISM: Mouse  
 39 <220> FEATURE:  
 40 <221> NAME/KEY: Unsure  
 41 <222> LOCATION: 5  
 42 <223> OTHER INFORMATION: Xaa is Ile or Val  
 43 <400> SEQUENCE: 2

W--> 44 Asn Trp Gly Arg Xaa Val Ala Phe Phe  
 45

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Output Set: N:\CRF4\01212003\I155327F.raw

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47 <210> SEQ ID NO: 3
48 <211> LENGTH: 31
49 <212> TYPE: DNA
50 <213> ORGANISM: Mouse
51 <220> FEATURE:
52 <221> NAME/KEY: modified_base
53 <222> LOCATION: 14
54 <223> OTHER INFORMATION: n is inosine
55 <220> FEATURE:
56 <221> NAME/KEY: modified_base
57 <222> LOCATION: 17
58 <223> OTHER INFORMATION: n is inosine
59 <220> FEATURE:
60 <221> NAME/KEY: modified_base
61 <222> LOCATION: 20
62 <223> OTHER INFORMATION: n is inosine
63 <400> SEQUENCE: 3
W--> 64 ggaattccca gccncntkn tcttggatcc a 31
66 <210> SEQ ID NO: 4
67 <211> LENGTH: 8
68 <212> TYPE: PRT
69 <213> ORGANISM: Mouse
70 <220> FEATURE:
71 <221> NAME/KEY: Unsure
72 <222> LOCATION: 4
73 <223> OTHER INFORMATION: Xaa is Asp or Glu
74 <220> FEATURE:
75 <221> NAME/KEY: Unsure
76 <222> LOCATION: 5
77 <223> OTHER INFORMATION: Xaa is Asn or Gln
78 <400> SEQUENCE: 4
W--> 79 Trp Ile Gln Xaa Xaa Gly Gly Trp
80 1 5
82 <210> SEQ ID NO: 5
83 <211> LENGTH: 14
84 <212> TYPE: PRT
85 <213> ORGANISM: Mouse
86 <400> SEQUENCE: 5
87 Met Ala Thr Pro Ala Ser Thr Pro Asp Thr Arg Ala Leu Val
88 1 5 10
90 <210> SEQ ID NO: 6
91 <211> LENGTH: 583
92 <212> TYPE: DNA
93 <213> ORGANISM: HUMAN
94 <220> FEATURE:
95 <221> NAME/KEY: CDS
96 <222> LOCATION: (1)..(579)
97 <400> SEQUENCE: 6
98 atg gcg acc cca gcc tcg gcc cca gac aca cg gct ctg gtg gca gac 48

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Input Set : N:\Crf4\01172003\I155327G.raw  
Output Set: N:\CRF4\01212003\I155327F.raw

99	Met Ala Thr Pro Ala Ser Ala Pro Asp Thr Arg Ala Leu Val Ala Asp	
100	1 5 10 15	
101	ttt gta ggt tat aag ctg agg cag aag ggt tat gtc tgt gga gct ggc	96
102	Phe Val Gly Tyr Lys Leu Arg Gln Lys Gly Tyr Val Cys Gly Ala Gly	
103	20 25 30	
104	ccc ggg gag ggc cca gca gct gac ccg ctg cac caa gcc atg cg gca	144
105	Pro Gly Glu Gly Pro Ala Ala Asp Pro Leu His Gln Ala Met Arg Ala	
106	35 40 45	
107	gct gga gat gag ttc gag acc cgc ttc cgg cgc acc ttc tct gat ctg	192
108	Ala Gly Asp Glu Phe Glu Thr Arg Phe Arg Arg Thr Phe Ser Asp Leu	
109	50 55 60	
110	gcg gct cag ctg cat gtg acc cca ggc tca gcc cag caa cgc ttc acc	240
111	Ala Ala Gln Leu His Val Thr Pro Gly Ser Ala Gln Gln Arg Phe Thr	
112	65 70 75 80	
113	cag gtc tcc gac gaa ctt ttt caa ggg ggc ccc aac tgg ggc cgc ctt	288
114	Gln Val Ser Asp Glu Leu Phe Gln Gly Pro Asn Trp Gly Arg Leu	
115	85 90 95	
116	gta gcc ttc ttt gtc ttt ggg gct gca ctg tgt gct gag agt gtc aac	336
117	Val Ala Phe Phe Val Phe Gly Ala Ala Leu Cys Ala Glu Ser Val Asn	
118	100 105 110	
119	aag gag atg gaa cca ctg gtg gga caa gtg cag gag tgg atg gtg gcc	384
120	Lys Glu Met Glu Pro Leu Val Gly Gln Val Glu Trp Met Val Ala	
121	115 120 125	
122	tac ctg gag acg cgg ctg gct gac tgg atc cac agc agt ggg ggc tgg	432
123	Tyr Leu Glu Thr Arg Leu Ala Asp Trp Ile His Ser Ser Gly Gly Trp	
124	130 135 140	
125	gcg gag ttc aca gct cta tac ggg gac ggg gcc ctg gag gag gcg cgg	480
126	Ala Glu Phe Thr Ala Leu Tyr Gly Asp Gly Ala Leu Glu Ala Arg	
127	145 150 155 160	
128	cgt ctg cgg gag ggg aac tgg gca tca gtg agg aca gtg ctg acg ggg	528
129	Arg Leu Arg Glu Gly Asn Trp Ala Ser Val Arg Thr Val Leu Thr Gly	
130	165 170 175	
131	gcc gtg gca ctg ggg gcc ctg gta act gta ggg gcc ttt ttt gct agc	576
132	Ala Val Ala Leu Gly Ala Leu Val Thr Val Gly Ala Phe Phe Ala Ser	
133	180 185 190	
134	aag tgaa	583
135	Lys	
137	<210> SEQ ID NO: 7	
138	<211> LENGTH: 193	
139	<212> TYPE: PRT	
140	<213> ORGANISM: HUMAN	
141	<400> SEQUENCE: 7	
142	Met Ala Thr Pro Ala Ser Ala Pro Asp Thr Arg Ala Leu Val Ala Asp	
143	1 5 10 15	
144	Phe Val Gly Tyr Lys Leu Arg Gln Lys Gly Tyr Val Cys Gly Ala Gly	
145	20 25 30	
146	Pro Gly Glu Gly Pro Ala Ala Asp Pro Leu His Gln Ala Met Arg Ala	
147	35 40 45	
148	Ala Gly Asp Glu Phe Glu Thr Arg Phe Arg Arg Thr Phe Ser Asp Leu	

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Input Set : N:\Crf4\01172003\I155327G.raw  
Output Set: N:\CRF4\01212003\I155327F.raw

149	50	55	60	
150	Ala Ala Gln Leu His Val Thr Pro Gly Ser Ala Gln Gln Arg Phe Thr			80
151	65	70	75	
152	Gln Val Ser Asp Glu Leu Phe Gln Gly Gly Pro Asn Trp Gly Arg Leu			95
153	85	90		
154	Val Ala Phe Phe Val Phe Gly Ala Ala Leu Cys Ala Glu Ser Val Asn			
155	100	105	110	
156	Lys Glu Met Glu Pro Leu Val Gly Gln Val Gln Glu Trp Met Val Ala			
157	115	120	125	
158	Tyr Leu Glu Thr Arg Leu Ala Asp Trp Ile His Ser Ser Gly Gly Trp			
159	130	135	140	
160	Ala Glu Phe Thr Ala Leu Tyr Gly Asp Gly Ala Leu Glu Glu Ala Arg			160
161	145	150	155	
162	Arg Leu Arg Glu Gly Asn Trp Ala Ser Val Arg Thr Val Leu Thr Gly			
163	165	170	175	
164	Ala Val Ala Leu Gly Ala Leu Val Thr Val Gly Ala Phe Phe Ala Ser			
165	180	185	190	
166	Lys			
168	<210> SEQ ID NO: 8			
169	<211> LENGTH: 582			
170	<212> TYPE: DNA			
171	<213> ORGANISM: Mouse			
172	<220> FEATURE:			
173	<221> NAME/KEY: CDS			
174	<222> LOCATION: (1)..(579)			
175	<400> SEQUENCE: 8			
176	atg gcg acc cca gcc tca acc cca gac aca cg <sup>48</sup>			
177	Met Ala Thr Pro Ala Ser Thr Pro Asp Thr Arg Ala Leu Val Ala Asp			
178	1 5 10 15			
179	ttt gta ggc tat agg ctg agg cag aag ggt tat gtc tgt gga gct ggc <sup>96</sup>			
180	Phe Val Gly Tyr Arg Leu Arg Gln Lys Gly Tyr Val Cys Gly Ala Gly			
181	20 25 30			
182	cct ggg gaa ggc cca gcc gac ccg ctg cac caa gcc atg cg <sup>144</sup>			
183	Pro Gly Glu Gly Pro Ala Ala Asp Pro Leu His Gln Ala Met Arg Ala			
184	35 40 45			
185	gct gga gac gag ttt gag acc cgt ttc cgc acc ttc tct gac ctg <sup>192</sup>			
186	Ala Gly Asp Glu Phe Glu Thr Arg Phe Arg Arg Thr Phe Ser Asp Leu			
187	50 55 60			
188	gcc gct cag cta cac gtg acc cca ggc tca gcc cag caa cgc ttc acc <sup>240</sup>			
189	Ala Ala Gln Leu His Val Thr Pro Gly Ser Ala Gln Gln Arg Phe Thr			
190	65 70 75 80			
191	cag gtt tcc gac gaa ctt ttc caa ggg ggc cct aac tgg ggc cgt ctt <sup>288</sup>			
192	Gln Val Ser Asp Glu Leu Phe Gln Gly Pro Asn Trp Gly Arg Leu			
193	85 90 95			
194	gtg gca ttc ttt gtc ttt ggg gct gcc ctg tgt gct gag agt gtc aac <sup>336</sup>			
195	Val Ala Phe Phe Val Phe Gly Ala Ala Leu Cys Ala Glu Ser Val Asn			
196	100 105 110			
197	aaa gaa atg gag cct ttg gtg gga caa gtg cag gat tgg atg gtg gcc <sup>384</sup>			
198	Lys Glu Met Glu Pro Leu Val Gly Gln Val Gln Asp Trp Met Val Ala			

**RAW SEQUENCE LISTING**  
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Input Set : N:\Crf4\01172003\I155327G.raw  
 Output Set: N:\CRF4\01212003\I155327F.raw

199	115	120	125	
200	tac ctg gag aca cgt ctg gct gac tgg atc cac agc agt ggc ggc tgg			432
201	Tyr Leu Glu Thr Arg Leu Ala Asp Trp Ile His Ser Ser Gly Gly Trp			
202	130	135	140	
203	gcg gag ttc aca gct cta tac ggg gac ggg gcc ctg gag gag gca cg			480
204	Ala Glu Phe Thr Ala Leu Tyr Gly Asp Gly Ala Leu Glu Ala Arg			
205	145	150	155	160
206	cgt ctg cgg gag ggg aac tgg gca tca gtg agg aca gtg ctg acg ggg			528
207	Arg Leu Arg Glu Gly Asn Trp Ala Ser Val Arg Thr Val Leu Thr Gly			
208	165	170	175	
209	gcc gtg gca ctg ggg gcc ctg gta act gta ggg gcc ttt ttt gct agc			576
210	Ala Val Ala Leu Gly Ala Leu Val Thr Val Gly Ala Phe Phe Ala Ser			
211	180	185	190	
212	aag tga			582
213	Lys			
215 <210>	SEQ ID NO: 9			
216 <211>	LENGTH: 193			
217 <212>	TYPE: PRT			
218 <213>	ORGANISM: Mouse			
219 <400>	SEQUENCE: 9			
220	Met Ala Thr Pro Ala Ser Thr Pro Asp Thr Arg Ala Leu Val Ala Asp			
221	1	5	10	15
222	Phe Val Gly Tyr Arg Leu Arg Gln Lys Gly Tyr Val Cys Gly Ala Gly			
223	20	25	30	
224	Pro Gly Glu Gly Pro Ala Ala Asp Pro Leu His Gln Ala Met Arg Ala			
225	35	40	45	
226	Ala Gly Asp Glu Phe Glu Thr Arg Phe Arg Arg Thr Phe Ser Asp Leu			
227	50	55	60	
228	Ala Ala Gln Leu His Val Thr Pro Gly Ser Ala Gln Gln Arg Phe Thr			
229	65	70	75	80
230	Gln Val Ser Asp Glu Leu Phe Gln Gly Gly Pro Asn Trp Gly Arg Leu			
231	85	90	95	
232	Val Ala Phe Phe Val Phe Gly Ala Ala Leu Cys Ala Glu Ser Val Asn			
233	100	105	110	
234	Lys Glu Met Glu Pro Leu Val Gly Gln Val Gln Asp Trp Met Val Ala			
235	115	120	125	
236	Tyr Leu Glu Thr Arg Leu Ala Asp Trp Ile His Ser Ser Gly Gly Trp			
237	130	135	140	
238	Ala Glu Phe Thr Ala Leu Tyr Gly Asp Gly Ala Leu Glu Ala Arg			
239	145	150	155	160
240	Arg Leu Arg Glu Gly Asn Trp Ala Ser Val Arg Thr Val Leu Thr Gly			
241	165	170	175	
242	Ala Val Ala Leu Gly Ala Leu Val Thr Val Gly Ala Phe Phe Ala Ser			
243	180	185	190	
244	Lys			
246 <210>	SEQ ID NO: 10			
247 <211>	LENGTH: 333			
248 <212>	TYPE: PRT			
249 <213>	ORGANISM: murine			

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/155,327F

DATE: 01/21/2003  
TIME: 12:26:17

Input Set : N:\Crf4\01172003\I155327G.raw  
Output Set: N:\CRF4\01212003\I155327F.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 16,19,22,25

Seq#:2; Xaa Pos. 5

Seq#:3; N Pos. 14,17,20

Seq#:4; Xaa Pos. 4,5

**VERIFICATION SUMMARY**

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Input Set : N:\Crf4\01172003\I155327G.raw  
Output Set: N:\CRF4\01212003\I155327F.raw

L:5 M:270 C: Current Application Number differs, Wrong Format  
L:6 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:33 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0